RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/581,228
Source: IFW
Date Processed by STIC: 6/9/06

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IFWP

RAW SEQUENCE LISTING DATE: 06/09/2006
PATENT APPLICATION: **US/10/581,228** TIME: 10:39:52

Input Set : **A:\140140417U2.SEQ.TXT**
Output Set: **N:\CRF4\06092006\J581228.raw**

<110> APPLICANT: John A. Chiorini
Michael Schmidt
Ioannis Bossis
Di Giovanni Pasquale
<120> TITLE OF INVENTION: Bovine Adeno-Associated Viral (BAAV)
Vector and Uses Thereof
<130> FILE REFERENCE: 14014.0417U2
C--> **<140> CURRENT APPLICATION NUMBER: US/10/581,228**
C--> **<141> CURRENT FILING DATE: 2006-06-02**
<150> PRIOR APPLICATION NUMBER: PCT/US04/40825
<151> PRIOR FILING DATE: 2004-12-06
<150> PRIOR APPLICATION NUMBER: 60/526,786
<151> PRIOR FILING DATE: 2003-12-04
<160> NUMBER OF SEQ ID NOS: 23
<170> SOFTWARE: FastSEQ for Windows Version 4.0
<210> SEQ ID NO: 1
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<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence; note=synthetic construct
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gattcaggag aatcaggaga gctacctgtc cttcaactcc acgggaaact ctcggagcca 1140
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58 gaactttccc tttaatgact gtgtggaaaa aatgttgatc tggtgggagg agggaaagat 1500
59 gaccagcaag gtggtggaac ccgccaaggc catcttgggg gggtctagag tacgagtgga 1560
60 tcaaaaatgt aaatcctctg tacaagtaga ctctaccccg gtgattatca cctccaatac 1620
61 taacatgtgt gtggtggtgg atgggaactc cacgaccttt gaacaccagc agccgctgga 1680
62 agaccgcatg ttcagatttg aactcatgcg gcggctcccg ccagattttg gcaagattac 1740
63 caagcaggaa gtcaaagact tttttgcttg ggcaaaggtc aaccaggtgc cggtgactca 1800
64 cgagtttatg gttcccaaga aagtggcggg aactgagagg gcggagactt ctagaaaacg 1860
65 cccactggat gacgtcacca ataccaacta taaaagtccg gagaagcggg cccggctctc 1920
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67 tctcaactgg tcttccaggt atgaatgcag atgtgactat catgctaaat ttgactctgt 2040
68 aacgggggaa tgtgacgagt gtgaatattt gaatcggggc aaaaatggct gtatctttca 2100
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70 agattttaat gattttgatg actgtaataa agagcagtaa ataaagtgag tagtcatgtc 2220
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83 ctgggtcctg ccgacctaca acaaccacct gtacctgcgg ctcggctcga gcaacgccag 3000
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89 gttcatggtg cctcagtacg ggtactgcgg actggtaacc ggaggcagct ctcaaaacca 3360
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91 aaacaacttt gagatggtgt acaagtttga aaacgtgccc ttccactcca tgtacgctca 3480
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94 gaccaaaaca aacttttctg gctaccgcaa aaactggctc ccggggccca tgatgaagca 3660
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98 tgcggggccc aacatcaccg gcaacaccac cacagatgcc aataacctga tgttcacttc 3900
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105 aatagaatgg gaaatccaga aggagcggtc caagagatgg aacccagagg tccagttcac 4320
106 gtccaactac ggagcacagg actcgcttct ctgggctccc gacaacgccg gagcctacaa 4380
107 agagcccagg gccattggat cccgatacct caccaaccac ctctagccca attctgttgc 4440
108 ataccctcaa taaaccgtgt attcgtttca gtaaaatact gcctcttgtg gtcattcggc 4500
109 gtacaacagc ttacaacaac aacaaaaccc ccttgctaga gagtgtggca ctccccccc 4560
110 tgtcgcgttc gctcgttcgc tggctcgatt gggggggtgg cagctcaaag agctgccaga 4620
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116 <212> TYPE: DNA
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120 <223> OTHER INFORMATION: Description of Artificial Sequence; note=synthetic
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123 <400> SEQUENCE: 2
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126 tcagatttga atttgactct gattgagcag cctcagctga cggtggctga cagaattcgc 180
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130 cagaacattg agccgcggat taacgactgg gtcgccatca ccaaggtaaa gaagggcgga 420
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133 gagcgcaaac ggctcgtcgc tcagtttcag cttgagtcct cgcagcgctc gcaagaggca 600
134 tcttcccaga gggacgtttc ggctgacccg gtcatcaaga gcaagacttc ccagaaatac 660
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Input Set : A:\140140417U2.SEQ.TXT
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170 Gln Ile Trp Glu Leu Pro Pro Glu Ser Asp Leu Asn Leu Thr Leu Ile
171 35 40 45
172 Glu Gln Pro Gln Leu Thr Val Ala Asp Arg Ile Arg Arg Val Phe Leu
173 50 55 60
174 Tyr Glu Trp Asn Lys Phe Ser Lys Gln Glu Ser Lys Phe Phe Val Gln
175 65 70 75 80
176 Phe Glu Lys Gly Ser Glu Tyr Phe His Leu His Thr Leu Val Glu Thr
177 85 90 95
178 Ser Gly Ile Ser Ser Met Val Leu Gly Arg Tyr Val Ser Gln Ile Arg
179 100 105 110
180 Ala Gln Leu Val Lys Val Val Phe Gln Asn Ile Glu Pro Arg Ile Asn
181 115 120 125
182 Asp Trp Val Ala Ile Thr Lys Val Lys Lys Gly Gly Ala Asn Lys Val
183 130 135 140
184 Val Asp Ser Gly Tyr Ile Pro Ala Tyr Leu Leu Pro Lys Val Gln Pro
185 145 150 155 160
186 Glu Leu Gln Trp Ala Trp Thr Asn Leu Glu Glu Tyr Lys Leu Ala Ala
187 165 170 175
188 Leu Asn Leu Glu Glu Arg Lys Arg Leu Val Ala Gln Phe Gln Leu Glu
189 180 185 190
190 Ser Ser Gln Arg Ser Gln Glu Ala Ser Ser Gln Arg Asp Val Ser Ala
191 195 200 205
192 Asp Pro Val Ile Lys Ser Lys Thr Ser Gln Lys Tyr Met Ala Leu Val
193 210 215 220
194 Ser Trp Leu Val Glu His Gly Ile Thr Ser Glu Lys Gln Trp Ile Gln
195 225 230 235 240
196 Glu Asn Gln Glu Ser Tyr Leu Ser Phe Asn Ser Thr Gly Asn Ser Arg
197 245 250 255
198 Ser Gln Ile Lys Ala Ala Leu Asp Asn Ala Ser Lys Ile Met Ser Leu
199 260 265 270
200 Thr Lys Ser Ala Ser Asp Tyr Leu Val Gly Gln Thr Val Pro Glu Asp
201 275 280 285
202 Ile Ser Glu Asn Arg Ile Trp Gln Ile Phe Asp Leu Asn Gly Tyr Asp
203 290 295 300
204 Pro Ala Tyr Ala Gly Ser Val Leu Tyr Gly Trp Cys Thr Arg Ala Phe
205 305 310 315 320
206 Gly Lys Arg Asn Thr Val Trp Leu Tyr Gly Pro Ala Thr Thr Gly Lys

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211         355                 360                 365
212 Met Leu Ile Trp Trp Glu Glu Gly Lys Met Thr Ser Lys Val Val Glu
213     370                 375                 380
214 Pro Ala Lys Ala Ile Leu Gly Gly Ser Arg Val Arg Val Asp Gln Lys
215 385                 390                 395                 400
216 Cys Lys Ser Ser Val Gln Val Asp Ser Thr Pro Val Ile Ile Thr Ser
217                 405                 410                 415
218 Asn Thr Asn Met Cys Val Val Val Asp Gly Asn Ser Thr Thr Phe Glu
219             420                 425                 430
220 His Gln Gln Pro Leu Glu Asp Arg Met Phe Arg Phe Glu Leu Met Arg
221         435                 440                 445
222 Arg Leu Pro Pro Asp Phe Gly Lys Ile Thr Lys Gln Glu Val Lys Asp
223     450                 455                 460
224 Phe Phe Ala Trp Ala Lys Val Asn Gln Val Pro Val Thr His Glu Phe
225 465                 470                 475                 480
226 Met Val Pro Lys Lys Val Ala Gly Thr Glu Arg Ala Glu Thr Ser Arg
227                 485                 490                 495
228 Lys Arg Pro Leu Asp Asp Val Thr Asn Thr Asn Tyr Lys Ser Pro Glu
229             500                 505                 510
230 Lys Arg Ala Arg Leu Ser Val Val Pro Glu Thr Pro Arg Ser Ser Asp
231         515                 520                 525
232 Val Pro Val Glu Pro Ala Pro Leu Arg Pro Leu Asn Trp Ser Ser Arg
233     530                 535                 540
234 Tyr Glu Cys Arg Cys Asp Tyr His Ala Lys Phe Asp Ser Val Thr Gly
235 545                 550                 555                 560
236 Glu Cys Asp Glu Cys Glu Tyr Leu Asn Arg Gly Lys Asn Gly Cys Ile
237                 565                 570                 575
238 Phe His Asn Ala Thr His Cys Gln Ile Cys His Ala Val Pro Pro Trp
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247 <212> TYPE: DNA
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256 gagaatcagg agagctacct gtccttcaac tccacgggaa actctcggag ccagattaaa     120
257 gccgcgcttg acaacgcgtc aaaaattatg agtctgacca aatctgcctc agactatctc     180
258 gtgggacaga ctgttccaga ggacatttct gaaaacagaa tctggcagat ttttgatctc     240
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L:14 M:270 C: Current Application Number differs, Replaced Current Application No
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date